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## SEQUENCE LISTING

### GENERAL INFORMATION:

- (i) APPLICANT: Frayne, Elizabeth Gay
- (ii) TITLE OF INVENTION: Microbial Production of Nuclease Resistant DNA, RNA, and Oligo Mixtures

(iii) NUMBER OF SEQUENCES: 2

### (iv) CORRESPONDENCE ADDRESS:

- (A) ADDRESSEE: Frayne, Elizabeth Gay
- (B) STREET: 2027 Galvin Ln. #1
- (C) CITY: Diamond Bar
- (D) STATE: California
- (E) COUNTRY: USA
- (F) ZIP: 91765

### (v) COMPUTER READABLE FORM:

- (A) MEDIUM TYPE: Diskette, 3.50 inch 1.4 MB
- (B) COMPUTER: IBM
- (C) OPERATING SYSTEM: Windows XP
- (D) SOFTWARE: Microsoft Word

### (vi) CURRENT APPLICATION DATA:

- (A) APPLICATION NUMBER: US 10/007,489
- (B) FILING DATE: 05-DEC-2001
- (C) CLASSIFICATION:

### (vii) PRIOR APPLICATION DATA:

- (A) APPLICATION NUMBER: US 60/100,231
- (B) FILING DATE: 14-SEP-1998
- (A) APPLICATION NUMBER: US 60/152,371

(B) FILING DATE: 07-SEP-1999

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: (909)861-2111/860-7415

(2) INFORMATION FOR SEQ ID NO:1:

(i)SEQUENCE CHARACTERISTICS:

(A) LENGTH: 22 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: genomic DNA

(iii) HYPOTHETICAL: no

(iv) ANTI-SENSE: no

(vi) ORIGINAL SOURCE:

(A) ORGANISM: *Saccharomyces cerevisiae*

(B) STRAIN: YFLO39C genomic clone

(x) PUBLICATION INFORMATION:

(A) AUTHORS: Murakami Y., Naitou M., Hagiwara H., Shibata T., Ozawa M.,  
Sasanuma S.I., Sasanuma M., Tsuchiya Y., Soeda E., Yokoyama K., Yamazaki M.,  
Tashiro H., Eki T.

(B) TITLE: Analysis of the nucleotide sequence of chromosome VI from  
*Saccharomyces cerevisiae*.

(C) JOURNAL: Nature Genet.

(D) VOLUME: 10

(F) PAGES: 261-268

(G) DATE:1995

(K) RELEVANT RESIDUES IN SEQ ID NO: 1: 10-31

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

GAG GTT GCT GCT TTG GTT ATT G 22

(2) INFORMATION FOR SEQ ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 22 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: genomic DNA

(iii) HYPOTHETICAL: no

(iv) ANTI-SENSE: yes

(vi) ORIGINAL SOURCE:

- (A) ORGANISM: *Saccharomyces cerevisiae*
- (B) STRAIN: YFLO39C genomic clone

(x) PUBLICATION INFORMATION:

- (A) AUTHORS: Murakami Y., Naitou M., Hagiwara H., Shibata T., Ozawa M., Sasanuma S.I., Sasanuma M., Tsuchiya Y., Soeda E., Yokoyama K., Yamazaki M., Tashiro H., Eki T.
- (B) TITLE: Analysis of the nucleotide sequence of chromosome VI from *Saccharomyces cerevisiae*.
- (C) JOURNAL: Nature Genet.
- (D) VOLUME: 10
- (F) PAGES: 261-268
- (G) DATE: 1995

(K) RELEVANT RESIDUES IN SEQ ID NO: 2: 1097-1118

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

TT GTG GTG AAC GAT AGA TGG AC

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